

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
E9QP59	E9QP59_MOUSE	Mus musculus	NaN	26.066638	NaN	NaN	28135057;28528544	MAAATAAAPQQLSDEELFSQLRRY GLSPGPVTESTRPVYLKLLKLRREEE QQQQQQQQQQHRAGGRGNKTRN SNNNTATAMGGRPGSGDLAYLRS PAGLGRLSASAAEFPVAGGSGGAAA VPAAGSKVLLGFSSDESDEVEASPRE QAGGGGARRDRAALQYRGLRAPPAP PAAGEVTGGHPGERRRPHSWWGAR RPAGPEPQPPAAGSDGAAEDADEEL ADGEDRDPEAEPLWASRTVNGSRL LPYSSCREHYSDSEEEEEEGEEDGD VAPARQVLKDDSLARHRPRRSHSKP FSALTAKSGGSRQETSVQGGGALAM NDRAAAAAGSLDRSRNLEEAAPGG GGGGCGCCDPVDSIPRYRAGAKKLA PLLSPPSPDGDSTLESPTGPLLKTNN HIGGCAFVDSPLYANSPLPGATA AAAPGTLRINHANHTGSNHTYLKTA YGKPKLCEPEEELLQQFKREEVSPT GSFSAHYLSMFLLLAACLFLLGLT YLGMRTGVPEDEGLIKNPFDETFG KIQESEKNLLMSTLYKLHDLRLAQIAG DHECGSSQRMLSVQEAAYLKNL GPEYEDVFNTSLLWIFKNGKDVGIR CVGYGPEEDLTNITDVQFLQSTRPQ MPFWCRFRRAFITVTHRLLLLCLGV VLVCVALRYMRYRWTKEEEETROM YDMVVKIIDVLRSHNEACQETKDLQ PYMPLPHVRDSLIPQDRKKMKKV WDRVDFLAANESRVTRTERRVGG ADFLVWRWIQPSASCDKTLVIPSJV WQQQAFHLDRRNSPPNSLTPCLKIR NMFDPVMEIGDHWHLAQEAILEKC SDNDGIVHIAVDRNSREGCVYVKCL SPEYAGKAFKALHGSWFDGKLVTVK YLRDLRYHHRFPQALCNTPLKPPAN KHMNSLSHLRLRTGLANSQGSS	True	False	1.507	5.0	1.077	1.875	1.359	3.366	1.258